

-continued

Lys	Thr	Ile	Ser	Lys	Ala	Lys	Gly	Gln	Pro	Arg	Glu	Pro	Gln	Val	Tyr
340															
															350
Thr	Leu	Pro	Pro	Ser	Arg	Asp	Glu	Leu	Thr	Lys	Asn	Gln	Val	Ser	Leu
355															
															365
Thr	Cys	Leu	Val	Lys	Gly	Phe	Tyr	Pro	Ser	Asp	Ile	Ala	Val	Glu	Trp
370															
															380
Glu	Ser	Asn	Gly	Gln	Pro	Glu	Asn	Asn	Tyr	Lys	Thr	Thr	Pro	Pro	Val
385															
															400
Leu	Asp	Ser	Asp	Gly	Ser	Phe	Phe	Leu	Tyr	Ser	Lys	Leu	Thr	Val	Asp
405															
															415
Lys	Ser	Arg	Trp	Gln	Gln	Gly	Asn	Val	Phe	Ser	Cys	Ser	Val	Met	His
420															
															430
Glu	Ala	Leu	His	Asn	His	Tyr	Thr	Gln	Lys	Ser	Leu	Ser	Leu	Ser	Pro
435															
															445
Gly	Lys														
450															

1. An anti-CD96 antibody comprising: (i) a first light chain hypervariable region (HVR-L1), a second light chain hypervariable region (HVR-L2), and a third light chain hypervariable region (HVR-L3), and/or (ii) a first heavy chain hypervariable region (HVR-H1), a second heavy chain hypervariable region (HVR-H2), and a third heavy chain hypervariable region (HVR-H3); wherein:

- (a) HVR-L1 comprises an amino acid sequence selected from KASQNVGTAIV (SEQ ID NO: 13), KSSQSLLDSDGKTYLN (SEQ ID NO: 17), RVSQD-ISFWLS (SEQ ID NO: 21), RASSNVKYM (SEQ ID NO: 25), KASQSVTFADTGLMH (SEQ ID NO: 29), RSSTGAVITTSNYAN (SEQ ID NO: 33), RASQDI-YRNLH (SEQ ID NO: 37), or RASQXIXXNXH (SEQ ID NO: 308), wherein X at position 5 is D, A, E, G, H, K, N, P, Q, S, or T; X at position 7 is Y, or F; X at position 8 is R, K, or Q; X at position 10 is L, I, M, or V;
- (b) HVR-L2 comprises an amino acid sequence selected from SASTRYT (SEQ ID NO: 14), LVSKLDS (SEQ ID NO: 18), KASNLT (SEQ ID NO: 22), YTSLNLAS (SEQ ID NO: 26), RASNLEV (SEQ ID NO: 30), GTNNRAP (SEQ ID NO: 34), HASDSIS (SEQ ID NO: 38), or HAXXXXS (SEQ ID NO: 325), wherein X at position 3 is S, or E; X at position 4 is D, E, K, or Q; X at position 5 is S, H, L, R, or V; X at position 6 is I, or V;
- (c) HVR-L3 comprises an amino acid sequence selected from QQYSSSPLT (SEQ ID NO: 15), LQATHSPQT (SEQ ID NO: 19), LQSQSYPYT (SEQ ID NO: 23), QQFTSSPLT (SEQ ID NO: 27), QQSREYPWT (SEQ ID NO: 31), SLWYGSHWV (SEQ ID NO: 35), LQGYSMPYT (SEQ ID NO: 39), or XQGYXMPXT (SEQ ID NO: 335), wherein X at position 1 is L, G, M, or Q; X at position 5 is S, A, E, Q, or V; X at position 8 is Y, or F;
- (d) HVR-H1 comprises an amino acid sequence selected from TNWNWMH (SEQ ID NO: 41), GYGVT (SEQ ID NO: 45), TDYYIN (SEQ ID NO: 49), NDYYIN (SEQ ID NO: 53), SDYYMY (SEQ ID NO: 57), TNYGIH (SEQ ID NO: 61), TTYGMS (SEQ ID NO: 65),

XNXXXXH (SEQ ID NO: 72), wherein X at position 1 is T, A, D, E, G, H, K, N, Q, R, S, V, W, or Y; X at position 3 is N, A, F, G, H, M, R, S, V, or Y; X at position 4 is W, or F; X at position 5 is M, A, D, E, F, G, L, N, Q, R, S, T, V, or W, or XXXGXS (SEQ ID NO: 344), wherein X at position 1 is T, A, D, E, G, H, K, M, N, Q, R, or S; X at position 2 is T, D, E, G, H, N, Q, or S; X at position 3 is Y, F, M, or Q; X at position 5 is M, I, L, or V;

(e) HVR-H2 comprises an amino acid sequence selected from MIHPNSGITNINE (SEQ ID NO: 42), EIYPGT-VITYYNA (SEQ ID NO: 46), WIFPGTEGIYYNE (SEQ ID NO: 50), WIFPGRIITYYNE (SEQ ID NO: 54), AISDDGTYTYYPD (SEQ ID NO: 58), IIWAGGSTNYNSA (SEQ ID NO: 62), WINTDS-GVPTYAD (SEQ ID NO: 66), XXHXXXXXXXXXXN (SEQ ID NO: 107), wherein X at position 1 is M or F; X at position 2 is I, L, M, or V; X at position 4 is P, A, D, E, F, G, H, I, K, L, M, N, Q, R, S, T, V, or W; X at position 5 is N, A, D, E, F, G, H, I, K, L, M, Q, R, S, T, V, W, or Y; X at position 6 is S, A, G, T, or V; X at position 7 is G, A, or S; X at position 8 is I, A, or V; X at position 9 is T, A, D, E, G, H, I, K, L, M, N, Q, R, S, V, W, or Y; X at position 10 is N, A, M, or S; X at position 11 is I, F, G, H, K, L, M, N, Q, R, S, T, V, W, or Y; X at position 13 is E, A, D, G, H, K, L, M, N, P, Q, R, S, T, V, W, or Y, or WINTXXGVPTYAD (SEQ ID NO: 369), wherein X at position 5 is D, or E; X at position 6 is S, or T;

(f) HVR-H3 comprises an amino acid sequence selected from RSDGTYEGYFDY (SEQ ID NO: 43), ARGL-GRAMDY (SEQ ID NO: 47), AREGDYRYYSPLY (SEQ ID NO: 51), ARGVGEGLFDY (SEQ ID NO: 55), AKAGSYDYFDV (SEQ ID NO: 59), ARVSMMG-FAY (SEQ ID NO: 63), ARNIYYGWNFDY (SEQ ID NO: 67), RXDXXXXXY (SEQ ID NO: 203), wherein X at position 2 is S, A, F, G, I, L, M, N, R, T, V, W, or Y; X at position 4 is G, or W; X at position 5 is T, D, E, F, H, I, K, L, M, N, Q, V, W, or Y; X at position 6 is Y, D, F, H, N, R, or W; X at position 7 is E, D, G, H,